

## SEQUENCE LISTING

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<120> Double-Stranded RNA-Mediated Gene Suppression

<130> J&J 2084

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<151> 2000-12-28

<150> AU PR3028

<151> 2001-02-09

<160> 35

<170> PatentIn version 3.1

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<213> Artificial Sequence

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34

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tgaggattca caaaccacaa ctagaatgca gtg

33

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<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> PCR Primer

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24

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acaaaccaca actagaatgc agtg

24

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29

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<400> 6  
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32

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<400> 7  
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34

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&lt;220&gt;

&lt;223&gt; PCR primer

&lt;400&gt; 8

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33

&lt;210&gt; 9

&lt;211&gt; 30

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;223&gt; PCR primer

&lt;400&gt; 9

tgaactagtt ctcggccgca tattaagtgc

30

&lt;210&gt; 10

&lt;211&gt; 24

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;223&gt; PCR primer

&lt;400&gt; 10

tgaaagctta agtttaaacg ctag

24

&lt;210&gt; 11

&lt;211&gt; 31

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;223&gt; PCR primer

&lt;400&gt; 11

gcgcactagt cgtattaccg ccatgcatta g

31

&lt;210&gt; 12

&lt;211&gt; 31

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;223&gt; PCR primer

&lt;400&gt; 12

gcgcactagt acgccttaag atacattgat g

31

&lt;210&gt; 13

&lt;211&gt; 31

<212> DNA  
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31

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<400> 14  
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<400> 15  
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33

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31

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<400> 19  
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31

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<400> 20  
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<400> 21  
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33

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<400> 22  
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33

<210> 23  
<211> 97  
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<400> 23  
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tgagtcctgt aggacgaaac atgcataggg ccctgat 97

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<220>  
<223> Oligonucleotide

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gacatgcatg gccctgataa gggccgaatt g 91

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<220>  
<223> PCR primer

<400> 25  
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<210> 26  
<211> 18  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> PCR primer

<400> 26  
gaacttgtag ccgtttac 18

<210> 27

<211> 21  
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<220>  
 <223> PCR primer

<400> 27  
 cgcagatcct gagcttgtat g 21

<210> 28  
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 <212> DNA  
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<220>  
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<400> 28  
 cactgcattc tagttgtg 18

<210> 29  
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<220>  
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<400> 29  
 ttaaccagct gtgggggagag ggctg 25

<210> 30  
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 <212> DNA  
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<220>  
 <223> primer

<400> 30  
 agccagcgat gcaaagatgg tgttg 25

<210> 31  
 <211> 559  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> region of the HIV-1 genome

<400> 31  
 aagcttgccct tgagtgcctc aagtagtgtg tgcccgtctg ttgtgtgact ctggtaacta 60

gagatccctc agaccctttt agtcagtgtg gaaaatctct agcagtggcg cccgaacagg 120  
gacctgaaag cgaaagggaa accagaggag ctctctcgac gcaggactcg gcttgctgaa 180  
gcgcgcacgg caagaggcga ggggcggcga ctggtgagta cgccaaaaat tttgactagc 240  
ggaggctaga aggagagaga tgggtgcgag agcgtcagta ttaagcgggg gagaattaga 300  
tcgatgggaa aaaattcggg taaggccagg gggaaagaaa aaatataaat taaaacatat 360  
agtatgggca agcaggggagc tagaacgatt cgcagttaat cctggcctgt tagaaacatc 420  
agaaggctgt agacaaatac tgggacagct acaaccatcc cttcagacag gatcagaaga 480  
acttagatca ttatataata cagtagcaac cctctattgt gtgcatcaaa ggatagagat 540  
aaaagacacc aaggaagct 559

<210> 32  
<211> 252  
<212> DNA  
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<220>  
<223> IVS ribozyme

<400> 32  
agatctggca ctgagtaatt gctgcagatc gtcaaaagca ggagtccttg agtagtctct 60  
agcatacggg acctactcaa gctatgcac aagcttggtg ccgagctcgg atccactagt 120  
aacggccgcc agtgtgctgg aattcgccct taagggcgaa ttctgcagat atcaagcttt 180  
ctagagtatg ctagtaatga cgatctgcag caatctgatg agtccttgag gacgaaactc 240  
agtgccagat ct 252

<210> 33  
<211> 559  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> fragment with antisense orientation of IVS ribozyme

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agcttccttg gtgtctttta tctctatcct ttgatgcaca caatagaggg ttgctactgt 60  
attatataat gatctaagtt cttctgatcc tgtctgaagg gatggttgta gctgtcccag 120  
tatttgtcta cagccttctg atgtttctaa caggccagga ttaactgcga atcgttctag 180  
ctccctgctt gcccatacta tatgttttaa tttatatattt ttctttcccc ctggccttaa 240



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ccgaattttt tcccatcgat ctaattctcc cccgcttaat actgacgctc tcgcacccat    300
ctctctcctt ctagcctccg ctagtcaaaa tttttggcgt actcaccagt cgccgcccct    360
cgcctcttgc cgtgcgcgct tcagcaagcc gagtcctgcg tcgagagagc tcctctgggt    420
tccctttcgc tttcaggtcc ctgttcgggc gccactgcta gagattttcc aactgacta    480
aaagggctctg agggatctct agttaccaga gtcacacaac agacgggcac acactacttg    540
aagcactcaa ggcaagctt                                                    559

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<210> 34
<211> 98
<212> DNA
<213> Artificial Sequence

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<220>
<223> region containing HIV Tar sequence

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<400> 34
gggtctctct ggtagacca gatctgagcc tgggagctct ctggctaact aggggaaccca    60
ctgcttaagc ctcaataaag cttgccttga gtgcttca                                98

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<210> 35
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<212> PRT
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<220>
<223> HIV-1 Tat amino acid sequence

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<400> 35

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Met Glu Pro Val Asp Pro Arg Leu Glu Pro Trp Lys His Pro Gly Ser
1           5           10           15

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Gln Pro Lys Thr Ala Cys Thr Asn Cys Tyr Cys Lys Lys Cys Cys Phe
          20           25           30

```

```

His Cys Gln Val Cys Phe Ile Thr Lys Ala Leu Gly Ile Ser Tyr Gly
          35           40           45

```

```

Arg Lys Lys Arg Arg Gln Arg Arg Arg Pro Pro Gln Gly Ser Gln Thr
          50           55           60

```

```

His Gln Val Ser Leu Ser Lys Gln Pro Thr Ser Gln Ser Arg Gly Asp
65           70           75           80

```

